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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO. CONFIRMATION NO.	
10/532,014	04/20/2005	Magnus Ingelman-Sundberg	25401-40	8991
24256 DINSMORE &	7590 04/19/200 SHOHL, LLP	EXAMINER		
1900 CHEMED	CENTER	DAVIS, MINH TAM B		
255 EAST FIFTH STREET CINCINNATI, OH 45202			ART UNIT PAPER NUMBER	
,			1642	
SHORTENED STATUTORY	Y PERIOD OF RESPONSE	MAIL DATE	DELIVERY MODE	
31 DAYS 04/19/2007 PAPER			PER	

Please find below and/or attached an Office communication concerning this application or proceeding.

If NO period for reply is specified above, the maximum statutory period will apply and will expire 6 MONTHS from the mailing date of this communication.

	Application No.	Applicant(s)			
	10/532,014	INGELMAN-SUNDBERG ET AL.			
Office Action Summary	Examiner	Art Unit			
	MINH-TAM DAVIS	1642			
The MAILING DATE of this communication app Period for Reply	ears on the cover sheet with the c	orrespondence address			
A SHORTENED STATUTORY PERIOD FOR REPLY WHICHEVER IS LONGER, FROM THE MAILING DA  - Extensions of time may be available under the provisions of 37 CFR 1.13 after SIX (6) MONTHS from the mailing date of this communication.  - If NO period for reply is specified above, the maximum statutory period w  - Failure to reply within the set or extended period for reply will, by statute, Any reply received by the Office later than three months after the mailing earned patent term adjustment. See 37 CFR 1.704(b).	ATE OF THIS COMMUNICATION 16(a). In no event, however, may a reply be tim ill apply and will expire SIX (6) MONTHS from cause the application to become ABANDONE	l. ely filed the mailing date of this communication. D (35 U.S.C. § 133).			
Status					
<ol> <li>Responsive to communication(s) filed on 21 December 2a)</li> <li>This action is FINAL. 2b)</li> <li>Since this application is in condition for allowant closed in accordance with the practice under Exercise.</li> </ol>	action is non-final. ace except for formal matters, pro				
Disposition of Claims					
<ul> <li>4)  Claim(s) 1-10 is/are pending in the application.</li> <li>4a) Of the above claim(s) is/are withdraw</li> <li>5)  Claim(s) is/are allowed.</li> <li>6)  Claim(s) is/are rejected.</li> <li>7)  Claim(s) 1-10 is/are objected to.</li> <li>8)  Claim(s) are subject to restriction and/or</li> </ul>	·	o ig o			
Application Papers					
9) The specification is objected to by the Examiner 10) The drawing(s) filed on is/are: a) acce Applicant may not request that any objection to the d Replacement drawing sheet(s) including the correction 11) The oath or declaration is objected to by the Examiner	epted or b) objected to by the Edrawing(s) be held in abeyance. See on is required if the drawing(s) is obj	37 CFR 1.85(a). ected to. See 37 CFR 1.121(d).			
Priority under 35 U.S.C. § 119					
<ul> <li>12) Acknowledgment is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d) or (f).</li> <li>a) All b) Some * c) None of:</li> <li>1. Certified copies of the priority documents have been received.</li> <li>2. Certified copies of the priority documents have been received in Application No</li> <li>3. Copies of the certified copies of the priority documents have been received in this National Stage application from the International Bureau (PCT Rule 17.2(a)).</li> <li>* See the attached detailed Office action for a list of the certified copies not received.</li> </ul>					
Attachment(s)  Notice of References Cited (PTO-892)  Notice of Draftsperson's Patent Drawing Review (PTO-948)	4) Interview Summary ( Paper No(s)/Mail Da	te			
B) Information Disclosure Statement(s) (PTO/SB/08) Paper No(s)/Mail Date	5)	atent Application			

Application/Control Number: 10/532,014

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## **DETAILED ACTION**

## Election/Restrictions

Restriction is required under 35 U.S.C. 121 and 372.

This application contains the following inventions or groups of inventions which are not so linked as to form a single general inventive concept under PCT Rule 13.1.

In accordance with 37 CFR 1.499, applicant is required, in reply to this action, to elect a single invention to which the claims must be restricted.

**Group 1.** Claims 1-3, drawn to a binding agent or an antibody to CYP2W1 or SEQ ID NO:8.

Groups 2-4. Claim 4, drawn to a method for treating lung, colon, or ovarian tumor, using CYP2W1 protein. A method treating each cancer constitutes a single, distinct invention.

Groups 5-7. Claim 4, drawn to a method for treating lung, colon, or ovarian tumor, using CYP2W1 nucleic acid. A method treating each cancer constitutes a single, distinct invention.

**Group 8.** Claims 5-6, drawn to a method for screening agent that modulates CYP2W1 protein.

**Group 9.** Claims 5-6, 10, drawn to a method for screening agent that modulates CYP2W1 nucleic acid, or genes regulated by CYP2W1 promoter.

Groups 10-12. Claim 7, drawn to a method for treating lung, colon, or ovarian cancer, using a substance activated by CYP2W1 protein. A method treating each cancer constitutes a single, distinct invention.

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Groups 13-15. Claim 7, drawn to a method for treating lung, colon, or ovarian cancer, using an inducer of CYP2W1 protein. A method treating each cancer constitutes a single, distinct invention.

Groups 16-18. Claim 7, drawn to a method for treating lung, colon, or ovarian cancer, using binding agent of CYP2W1 protein. A method treating each cancer constitutes a single, distinct invention.

Groups 19-21. Claim 7, drawn to a method for treating lung, colon, or ovarian cancer, using a combination of a substance activated by CYP2W1 protein, and an inducer of CYP2W1 protein, or a combination of a substance activated by CYP2W1 protein, an inducer of CYP2W1 protein, and a binding agent for CYP2W1 protein. A method treating each cancer constitutes a single, distinct invention.

Group 22. Claims 8-9, drawn to a DNA molecule, SEQ ID NO:10.

The inventions are distinct, each from the other because of the following reasons:

According to PCT Rule 13.2, unity of invention exists only when the shared same or corresponding technical feature is a contribution over the prior art. The inventions listed as groups 1-22 do not relate to a single general inventive concept because they lack the same or corresponding special technical feature. The technical feature of group I, an antibody binding specifically to CYP2W1 protein, or SEQ ID NO:8, is shown to be the same as the antibody taught by WO 200290521-A2 (Becha et al), or WO 200259260 A2 (Asundi et al), as evidenced by Banki et al, 1994, JBC, 269 (4): 2847-51, or Bendayan et al, 1995, J Histochem Cytochem,

Protein

23. .464

43(9): 881-886. Thus the invention of group I lacks novelty and does not make a contribution over the prior art.

WO 200290521-A2 teaches an antibody to the disclosed polypeptide, which is a human drug metabolizing enzyme (items 6, 14). The polypeptide taught by WO 200290521-A2 is 87% similar to SEQ ID NO:8, from amino acid 1 to amino acid 431 (MPSRCH search result, 2007, us-10-532-014.rag, result 2, pages 1-2).

```
MPSRCH search result, 2007, us-10-532-014.8.rag.
RESULT 2
AAE33381
     AAE33381 standard; protein; 464 AA.
XX
AC
    AAE33381;
XX
DT
     02-APR-2003 (first entry)
XX
DE
     Human DME-7 protein.
XX
KW
     Human; drug metabolizing enzyme; DME; gastrointestinal disorder; asthma;
KW
    peptic oesophagitis; peptic ulcer; Crohn's disease; autoimmune disorder;
     liver disorder; acquired immune deficiency syndrome; cataract; anaemia;
KW
     inflammatory disorder; developmental disorder; achondroplastic dwarfism;
KW
     Cushing's syndrome; endocrine disorder; diabetes insipidus; leukaemia;
KW
     Sheehan syndrome; hypothyroidism; metabolic disorder; Addison's disease;
KW
     glycogen storage disease; hypocalcaemia; conjunctivitis; pancreatitis;
KW
     adenocarcinoma; cell proliferative disorder; actinic keratosis; cancer;
KW
     arteriosclerosis; gene therapy; protein replacement therapy; lymphoma;
KW
     eye disorder; sarcoma; obesity; melanoma; myeloma; AIDS; gout.
XX
OS
     Homo sapiens.
XX
FH
                     Location/Qualifiers
     Key
FT
     Peptide
                     1. .28
FT
                     /label= Signal peptide
FT
     Peptide
                     1. .24
FT
                     /label= Signal peptide
FT
     Peptide
                     1. .22
FT
                     /label= Signal peptide
FT
    Peptide
                     1. .20
FT
                     /label= Signal peptide
FT
    Domain
                     4. .21
                     /note= "Transmembrane domain"
FT
FT
    Protein
                     21. .464
FT
                     /note= "Mature human DME protein"
```

```
FT
                     /note= "Mature human DME protein"
FT
                     23. .464
     Protein
                     /note= "Mature human DME protein"
FT
FT
     Protein
                     /note= "Mature human DME protein"
FT
FT
     Domain
                     194. .222
                     /note= "Transmembrane domain"
FT
XX
PN
     WO200290521-A2.
XX
PD
     14-NOV-2002.
XX
PF
     10-MAY-2002; 2002WO-US015052.
XX
PR
     10-MAY-2001; 2001US-0290430P.
PR
     08-JUN-2001; 2001US-0296880P.
PR
     22-JUN-2001; 2001US-0300472P.
PR
     29-JUN-2001; 2001US-0301794P.
XX
PA
     (INCY-) INCYTE GENOMICS INC.
XX
PΤ
     Ring HZ, Hafalia AJA, Sanjanwala MM, Yao MG, Zebarjadian Y;
PΙ
     Edwards CM, Yue H, Tang YT, Lee EA, Emerling BM, Warren BA,
PG;
                Thangavelu K, Becha SD, Huang J, Ding L, Li JX;
PΙ
     Nguyen DB,
PΙ
     Griffin JA, Forsythe IJ, Richardson TW;
XX
DR
    WPI; 2003-111969/10.
DR
    N-PSDB; AAD51048.
XX
PT
     New human drug metabolizing enzyme proteins and polynucleotides, useful
PT
     for diagnosing, treating or preventing gastrointestinal (e.g. Crohn's
PT
     disease) or autoimmune/inflammatory disorders (e.g. AIDS),
hypothyroidism
PT
    or cancer.
XX
PS
     Claim 1; Page 158-159; 175pp; English.
XX
CC
     The invention relates to human drug metabolizing enzyme (DME) proteins
CC
     and nucleic acid molecules encoding such proteins. Sequences of the
     invention are useful for diagnosing, treating or preventing
CC
CC
     gastrointestinal disorders (e.g. peptic oesophagitis, peptic ulcer or
CC
     Crohn's disease) including liver disorders, autoimmune/inflammatory
CC
     disorders (e.g. acquired immune deficiency syndrome; AIDS, anaemia,
CC
     asthma, gout, pancreatitis or Crohn's disease), developmental disorders
CC
     (e.g. Cushing's syndrome or achondroplastic dwarfism), endocrine
CC
     disorders (e.g. Sheehan syndrome, diabetes insipidus or hypothyroidism),
CC
     eye disorders (e.g. conjunctivitis or cataract), metabolic disorders
CC
     (e.g. Addison's disease, glycogen storage diseases, hypocalcaemia or
CC
     obesity) or cell proliferative disorders (e.g. actinic keratosis,
CC
     arteriosclerosis or cancers including adenocarcinoma, leukaemia,
CC
     lymphoma, melanoma, myeloma, sarcoma or bone, ovary, lung, breast,
CC
     prostate or skin cancer). The present sequence is human DME protein
XX
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```
SQ
    Sequence 464 AA;
 Query Match
                    87.8%; Score 2249; DB 6; Length 464;
 Best Local Similarity
                    99.5%; Pred. No. 6.1e-211;
 Matches 429: Conservative
                         2;
                            Mismatches
                                       0;
                                          Indels
                                                    Gaps
0;
         1 MALLLLLFIGLIGLWGLLCACAQDPSPAARWPPGPRPLPLVGNLHLLRLSQQDRSLMELS 60
Qу
           Db
         1 MALLLLLFLGLLGLWGLLCACAODPSPAARWPPGPRPLPLVGNLHLLRLSOODRSLMELS 60
        61 ERYGPVFTVHLGRQKTVVLTGFEAVKEALAGPGQELADRPPIAIFOLIORGGGIFFSSGA
Qу
120
           Db
        61 ERYGPVFTVHLGROKTVVLTGFEAVKEALAGPGOELADRPPIAIFOLIORGGGIFFSSGA
120
Qу
       121 RWRAAROFTVRALHSLGVGREPVADKILOELKCLSGOLDGYRGRPFPLALLGWAPSNITF
180
           121 RWRAAROFTVRALHSLGVGREPVADKILOELKCLSGOLDGYRGRPFPLALLGWAPSNITF
Db
180
Qу
       181 ALLFGRRFDYRDPVFVSLLGLIDEVMVLLGSPGLQLFNVYPWLGALLQLHRPVLRKIEEV
240
       Db
240
       241 RAILRTLLEARR PHVCPGDPVCSYVDALI QOGOGDDPEGLFAEANAVACTLDMVMAGTET
Qу
300
           Db
       241 RAILRTLLEARRPHVCPGDPVCSYVDALIQQGQGDDPEGLFAEANAVACTLDMVMAGTET
300
Qу
       301 TSATLQWAALLMGRHPDVQGRVQEELDRVLGPGRTPRLEDQQALPYTSAVLHEVQRFITL
360
           Db
       301 TSATLQWAALLMGRHPDVQGRVQEELDRVLGPGRTPRLEDQQALPYTSAVLHEVQRFITL
360
Qу
       361 LPHVPRCTAADTQLGGFLLPKGTPVIPLLTSVLLDETOWOTPGOFNPGHFLDANGHFVKR
420
           Db
       361 LPHVPRCTAADTQLGGFLLPKGTPVIPLLTSVLLDETQWQTPGQFNPGHFLDANGHFVKR
420
       421 EAFLPFSAGRR 431
Qу
           | | | | | | | | | | | | | | | | |
       421 EAFLPFSAGQQ 431
Db
```

Page 6

XX

WO 200259260 A2 teaches an antibody to the disclosed polypeptide. The polypeptide taught by WO 200290521-A2 is 75% similar to SEQ ID NO:8, from amino acid 57 to amino acid 431 (MPSRCH search result, 2007, us-10-532-014.rag, result 5, pages 2-4).

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MPSRCH search result, 2007, us-10-532-014.8.rag
RESULT 5
ABP64996
ID
     ABP64996 standard; protein; 408 AA.
XX
AC
     ABP64996;
XX
DT
     25-FEB-2003 (first entry)
XX
DE
     Human protein SEQ ID 656.
XX
KW
     Human; expressed sequence tag; EST; haematopoietic disorder;
KW
     central nervous system disease; viral infection;
     peripheral nervous system disease; non-healing wound; infectious
disease;
     immune deficiency; immune disorder; bacterial infection; allergy;
cancer;
KW
     fungal infection; autoimmune disorder; coagulation disorder; nootropic;
KW
     antiallergic; antiinflammatory; immunosuppressive; neuroprotective;
KW
     cytostatic; haemostatic; virucide; antibacterial; fungicide;
     immunostimulant; cerebroprotective.
KW
XX
OS
     Homo sapiens.
XX
PN
     WO200259260-A2.
XX
PD
     01-AUG-2002.
XX
PF
     16-NOV-2001; 2001WO-US042950.
XX
     17-NOV-2000; 2000US-00714936.
PR
XX
PA
     (HYSE-) HYSEO INC.
XX
ΡI
     Tang YT, Goodrich RW, Liu C, Zhou P, Asundi V,
                                                          Zhang J,
ΡI
     Ren F, Xue AJ, Yang Y, Wehrman T, Drmanac RT;
XX
DR
     WPI; 2002-590824/63.
DR
     N-PSDB; ABQ99582.
XX
PT
     New isolated polynucleotide, useful in research, diagnostic or
PT
     therapeutic methods, e.g. preventing or treating disorders involving
PT
     aberrant protein expression or biological activity.
```

```
PS
    Claim 20; SEQ ID NO 656; 394pp; English.
XX
CC
    The present invention relates to novel human coding sequences (ABQ99268-
    ABQ99608) and proteins (ABP64682-ABP65022). The sequences are useful in
CC
    therapeutic, diagnostic and research methods. The polynucleotides may be
CC
CC
    used in the field of molecular biology as hybridisation probes, primers
    for PCR, for chromosome and gene mapping, for the recombinant production
CC
CC
    of protein, or in generation of anti-sense DNA or RNA. The
CC
    polynucleotides are useful in diagnostics as expressed sequence tags
CC
    (ESTs) for identifying expressed genes or for physical mapping of the
CC
    human genome. The proteins may be used as molecular weight markers, or .
as
CC
    nutritional sources or supplements. The proteins may be used to maintain
CC
    and expand cell population in a totipotential or pluripotential state
CC
    useful for re-engineering damaged or diseased tissues, transplantation,
    manufacture of bio-pharmaceuticals or the development of bio-sensors.
CC
The
CC
    polynucleotides and proteins are useful for preventing, treating or
CC
    ameliorating disorders involving aberrant protein expression or
CC
    biological activity, e.g. haematopoietic disorders, central/peripheral
    nervous system diseases, mechanical and traumatic disorders, non-healing
CC
    wounds, immune deficiencies and disorders, infectious diseases caused by
CC
CC
    viral, bacterial or fungal infection, autoimmune disorders, allergic
CC
    reactions and conditions, coagulation disorders, or cancer. The
CC
    polynucleotide sequences of the invention were assembled from ESTs
CC
    isolated mainly by sequencing by hybridisation, and in some cases,
CC
    sequences obtained from one or more public databases. Note: The sequence
    data for this patent did not form part of the printed specification, but
CC
CC
    was obtained in electronic format directly from WIPO at
CC
    ftp.wipo.int/pub/published pct sequences
XX
SO
    Sequence 408 AA;
 Query Match
                        75.9%; Score 1945; DB 5; Length 408;
 Best Local Similarity
                        99.2%; Pred. No. 3.2e-181;
 Matches 372; Conservative
                              3; Mismatches
                                                  Indels
                                                               Gaps
0;
Qу
          57 MELSERYGPVFTVHLGRQKTVVLTGFEAVKEALAGPGQELADRPPIAIFQLIQRGGGIFF
116
             Db
           1 MELSERYGPVFTVHLGRQKTVVLTGFEAVKEALAGPGQELADRPPIAIFQLIQRGGGIFF 60
Qу
         117 SSGARWRAARQFTVRALHSLGVGREPVADKILQELKCLSGQLDGYRGRPFPLALLGWAPS
176
             Db
          61 SSGARWRAARQFTVRALHSLGVGREPVADKILOELKCLSGOLDGYRGRPFPLALLGWAPS
120
Qу
         177 NITFALLFGRRFDYRDPVFVSLLGLIDEVMVLLGSPGLOLFNVYPWLGALLOLHRPVLRK
236
             121 NITFALLFGRRFDYRDPVFVSLLGLIDEVMVLLGSPGLQLFNVHPWLGALLQLHRPVLRK
Db
180
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Qy 296	237	IEEVRAILRTLLEARRPHVCPGDPVCSYVDALIQQGQGDDPEGLFAEANAVACTLDMVMA
Db 240	181	
Qy 356	297	GTETTSATLQWAALLMGRHPDVQGRVQEELDRVLGPGRTPRLEDQQALPYTSAVLHEVQR
Db 300	241	
Qy 416	357	FITLLPHVPRCTAADTQLGGFLLPKGTPVIPLLTSVLLDETQWQTPGQFNPGHFLDANGH
Db 360	301	
Qy	417	FVKREAFLPFSAGRR 431
Db	361	FVKREAFLPFSAGQQ 375

The antibody taught by the art would bind to SEQ ID NO:8, or CYP2W1, which is interpreted as a variant of SEQ ID NO:8, as evidenced by Banki et al, which teach that an antibody against human transaldolase could bind to yeast transaldolase which is about 58% homologous with human transaldolase, i.e. an antibody could cross-react and bind to a polypeptide at least with 58% homology to its antigen (abstract), or Bendayan et al, which teach that anti-human proinsulin monoclonal antibody to the Arg-Arg dipeptide, although providing very specific binding results, cross-reacts with non-related molecules, i.e., rat, bovine, porcine and human glucagons (abstract).

Applicant is advised that the reply to this requirement to be complete must include an election of the invention to be examined even though the requirement be traversed (37 CFR 1.143).

Applicant is reminded that upon the cancellation of claims to a non-elected

invention, the inventorship must be amended in compliance with 37 CFR 1.48(b) if one or more of the currently named inventors is no longer an inventor of at least one claim remaining in the application. Any amendment of inventorship must be accompanied by a petition under 37 CFR 1.48(b) and by the fee required under 37 CFR 1.17(i).

Any inquiry concerning this communication or earlier communications from the examiner should be directed to MINH-TAM DAVIS whose telephone number is 571-272-0830. The examiner can normally be reached on 9:00 AM-5:30 PM.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, SHANON FOLEY can be reached on 571-272-0898. The fax phone number for the organization where this application or proceeding is assigned is 571-273-8300.

Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see http://pair-direct.uspto.gov. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free). If you would like assistance from a USPTO Customer Service Representative or access to the automated information system, call 800-786-9199 (IN USA OR CANADA) or 571-272-1000.

MINH TAM DAVIS April 12, 2007